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Addendum

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In the paper "CRANK: New Methods for Automated Macromolecular Crystal Structure Solution" by Ness et al. (2004, Structure *12*, 1753–1761), crystallographic programs were mentioned, but not included in the reference list. These programs were ARP/wARP (Perrakis et al., 1999), REFMAC (Murshudov et al., 1997), DM (Cowtan, 1994), SHELXE (Sheldrick, 2002), and SOLOMON (Abrahams and Leslie, 1996).

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